

re-run



1600

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/470,667

DATE: 02/26/2003

TIME: 13:27:00

Input Set : N:\AMC\US09470667.raw

Output Set: N:\CRF4\02262003\I470667.raw

1 <110> APPLICANT: Asakura, Akira  
 2 Hoshino, Tatsuo  
 3 Ojima, Setsuko  
 4 Shinjoh, Masako  
 5 Tomiyama, Noribumi  
 6 <120> TITLE OF INVENTION: Novel Alcohol/Aldehyde Dehydrogenases  
 7 <130> FILE REFERENCE: C38435/109700CON  
 8 <140> CURRENT APPLICATION NUMBER: US/09/470,667  
 9 <141> CURRENT FILING DATE: 1999-12-22  
 10 <150> PRIOR APPLICATION NUMBER: US 08/934,506  
 11 <151> PRIOR FILING DATE: 1997-09-19  
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 13 <170> SOFTWARE: PatentIn version 3.1  
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 18 <213> ORGANISM: Gluconobacter oxydans  
 19 <400> SEQUENCE: 1

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57      tggatcagct acggtcagaa ccaagaaaac taccgtcact cgccctgac gcagatcacg 180
58      actgagaacg tcggccaact gcaactggtc tgggcgcgcg gcatgcagcc gggcaaagtc 240
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62      gtttactttg tttcgtggga caaccacctg gtcgcctctg acaccgcaac tggccaagtg 480
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92      atcaactacg gtcagaacca agagaactac cgccactcgc cctgacgca gattaccgca 180
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94      gtgacccgcg ttgtccatga cggcgctatg tatctggcaa acccggtga cgtgatccag 300
95      gccatcgacg ccgcgaccgg cgatctgatc tgggaacacc gccgccaact gccgaacatc 360

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161 &lt;222&gt; LOCATION: (1)..(23)

162 &lt;223&gt; OTHER INFORMATION:

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169 35 40 45
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171 50 55 60
172 Gly Gln Leu Gln Leu Val Trp Ala Arg Gly Met Gln Pro Gly Lys Val
173 65 70 75 80
174 Gln Val Thr Pro Leu Ile His Asp Gly Val Met Tyr Leu Ala Asn Pro
175 85 90 95
176 Gly Asp Val Ile Gln Ala Ile Asp Ala Lys Thr Gly Asp Leu Ile Trp
177 100 105 110
178 Glu His Arg Arg Gln Leu Pro Asn Ile Ala Thr Leu Asn Ser Phe Gly
179 115 120 125
180 Glu Pro Thr Arg Gly Met Ala Leu Tyr Gly Thr Asn Val Tyr Phe Val
181 130 135 140
182 Ser Trp Asp Asn His Leu Val Ala Leu Asp Thr Ala Thr Gly Gln Val
183 145 150 155 160
184 Thr Phe Asp Val Asp Arg Gly Gln Gly Glu Asp Met Val Ser Asn Ser
185 165 170 175
186 Ser Gly Pro Ile Val Ala Asn Gly Val Ile Val Ala Gly Ser Thr Cys
187 180 185 190
188 Gln Tyr Ser Pro Phe Gly Cys Phe Val Ser Gly His Asp Ser Ala Thr
189 195 200 205
190 Gly Glu Glu Leu Trp Arg Asn Tyr Phe Ile Pro Arg Ala Gly Glu Glu
191 210 215 220
192 Gly Asp Glu Thr Trp Gly Asn Asp Tyr Glu Ala Arg Trp Met Thr Gly
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201           290           295           300
202 Asp Gln Glu Cys Thr Phe Glu Met Met Val Thr Asn Val Asp Val Gln
203           305           310           315           320
204 Pro Ser Thr Glu Met Glu Gly Leu Gln Ser Ile Asn Pro Asn Ala Ala
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206 Thr Gly Glu Arg Arg Val Leu Thr Gly Val Pro Cys Lys Thr Gly Thr
207           340           345           350
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209           355           360           365
210 Thr Asn Tyr Gln Asn Met Ile Glu Ser Ile Asp Glu Asn Gly Ile Val
211           370           375           380
212 Thr Val Asn Glu Asp Ala Ile Leu Lys Glu Leu Asp Val Glu Tyr Asp
213           385           390           395           400
214 Val Cys Pro Thr Phe Leu Gly Gly Arg Asp Trp Pro Ser Ala Ala Leu
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216 Asn Pro Asp Ser Gly Ile Tyr Phe Ile Pro Leu Asn Asn Val Cys Tyr
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245 <223> OTHER INFORMATION:

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